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Result
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length: 2000000000
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Match Length
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14.250 Million cell updates/sec
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geneseqp1990s:*
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(c) 1993 - 2006 Biocceleration Ltd.
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rECEF 84.
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. !	Ado59801 B. subtil	Adk90667 Baker's y	Abg93115 S. cerevi	Abp60801 Gallus ga	Abp60810 Rattus no	Abp60809 Oryctolag	Abp60806 Mus muscu	Abp60805 Macaca mu	Abp60812 Ovis arie	Abp60797 Bos tauru	Abp60803 Homo sapi	Abp60927 Staphyloc	Aag80708 S. cerevi	Aar72389 Recombina	Aar47855 Human ADF	Aar42820 rECEF 104	Adm32933 Amino aci	Adk90669 Baker's y	Abr41607 Human DIT	Abp73948 Candida a	Abg93328 C. albica	

ALIGNMENTS

RESULT 1 AAP71087 ID AAP7 25-MAR-2003 31-OCT-2002 06-JUN-1991 Misc-difference Unidentified. Redox active peptide; antioxidant; stress; ischaemia; lipid peroxidation; food industry; cosmetics; antibiotic. Sequence of thioredoxin analogue. AAP71087; AAP71087 standard; peptide; 6 AA. (first entry) (revised) /note= "optional, and may optionally be joined to a terminal blocking gp." Location/Qualifiers

XZX 10-FEB-1986; 14-MAR-1986; 20-OCT-1986; WPI; 1987-258442/37. Pigiet VP, Millis CD; 10-FEB-1987; EP237189-A. Misc-difference (REPK) REPLIGEN CORP. 16-SEP-1987. 86US-00921287. 86US-00828112. 86US-00839857. 87EP-00301150 /note= "see above"

It is a redox active peptide which can be used in cpds. to treat stress

Claim 7; Page 7; 7pp; English

states.

New and known thioredoxin derivs. metal catalysed oxidative damage

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and analogues in biological :

s - useful for reactions and

preventing disease

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RESULT 2
ABP54936
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Best Local
                            dithiols. The results indicated that GSAO selectively binds proteins containing closely spaced thiols. The invention provides a method for the treatment and/or prophylaxis of arthritis using a compound of formula A-(L-Y)p, where A is a cell-membrane impermeable pendant group, L is a linker and/or spacer, Y is an arsenoxide or arsenoxide equivalent, and p is an integer from 1 to 10. The compound, e.g. GSAO, is useful for the treatment of arthritic conditions such as calcific periarthritis, enteropathic, chronic, thumb, hip, peripheral, psoriatic, rheumatoid, septic or Jaccoud's arthritis, gout, hand or knee osteoarthritis, oligoarthritis, and polyarthritis (all claimed). The compound blocks angiogenesis in the synovial tissue and leucocyte interest.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and ischaemia, for inhibiting lipid peroxidn. due to drugs or toxic effects, as anti-inflammatory agents, to prevent DNA damage, and to maintain the biological activity of antiblotics or proteins. Thioredoxin can be used as a general antioxidant in the food and cosmetic industry. It may be used in prodm. processes for antibiotics, proteins or polypeptides. Therapeutic dose is 1 microgram/kg - 100mg/kg. (Updated on 11-0CT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is that of a hexapeptide comprising the active site sequence of thioredoxin. The hexapeptide was used in an example from the invention that examined the binding of 4-(N-(S-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Use of an impermeable cell membrane compound for the treatment arthritis in vertebrates.  \label{eq:continuous} % \begin{subarray}{ll} \end{subarray} % \begin{subarray}{ll} \end{suba
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ingress that triggers
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Pred. No.
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no signs or symptoms
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Sequence 6

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1 WCGPCK

Query Match Best Local Matches

Similarity 6

100.0%;

Score 47; Pred. No. Mismatches

2e+06 DB 6;

Length 6;

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                                                                                     The present invention relates to a method for producing an antiangiogenesis plasmin fragment. The method comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin reductase, where a reduced plasmin protein is produced and the antiangiogenesis plasmin fragment having anti-angiogenesis activity, is released from the reduced plasmin protein. The plasminogen activator is preferably a urokinase-type plasminogen activator. The angiogenesis plasmin fragment is A61. The plasmin reductase is selected from annexin II heterotetramer, annexin II p36 subunit, p11, thioredoxin, and protein disulphide isomerase. The annexin II heterotetramer is associated with a cell membrane. The method of the invention is useful for modulating (e.g. promoting or inhibiting) angiogenesis. The present sequence represents a mammalian cleavage peptide found in thioredoxin and protein disulphide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein distlibhide isomerase; modulating ancienced plasmin protein; anti-angiogenesi urokinase-type plasminogen activator; A61; annexin II heterc annexin II p36 subunit; annexin II p11 subunit; thioredoxin; protein distlibhide isomerase; modulating angicanners.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Producing an anti-angiogenesis plasmin fragment, useful for modulating, e.g. promoting or inhibiting angiogenesis, comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammalian
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   Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-596985/56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Waisman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-NOV-2001; 2001US-0333866P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-NOV-2002; 2002US-00304287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cleavage peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (WAIS/) WAISMAN D. (KWON/) KWON M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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ilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide; 6
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Pred. No. 2e+06;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      anti-angiogenesis activity; annexin II heterotetramer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    plasminogen activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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Sequence 6 AA
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The present sequence represents an active site of a

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                                                                                                                                                                                                                                                                                                                                                                                                                                       a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic composition. The protein comprises thioredoxin selected from prokaryotic
                                                                                                                                                                                                                                                                           dinucleotide phosphate (reduced form) (NADPH) for reducing the thioredoxin active site of the protein, and thioredoxin reductase method and composition are useful for decreasing the viscosity and
                                                                                                                                                                                                                                                                                                                                           thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian thioredoxin. The composition further comprises nicotinamide-adenine dinucleotide phosphate (reduced form) (NADPH) for reducing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum. The method comprises contacting the mucus or sputum of the patient with a composition comprising a protein or peptide containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 13; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprising a protein or
reduced state.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-SEP-2002;
11-APR-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mucus; sputum;
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                                                                                                                                   cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NAJE-) NAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence of
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2003US-0462082P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 51; 69pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "this residue is in a reduced state"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "this residue is"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note= "active site of thioredoxin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the
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Matches 6
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Best Local
                                                                                                                                                                                                        The present invention described 4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2,5-dienone derivatives (I) and their salts, esters, amides, solvates, hydrates or protected forms. (I) have cytostatic activities, and can be used as thioredoxin reductase inhibitors. (I) can be used for treating a proliferative condition (especially colon or renal cancer) or a condition mediated by thioredoxin/thioredoxin reductase in the human or animal body. (I) are also useful for inhibiting thioredoxin/thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2; 5-dienone; cytostatic; thioredoxin reductase inhibitor; proliferative condition; colon cancer; renal cancer; thioredoxin; thioredoxin reductase; cell cycle progression; cell proliferation; apoptosis; catalytic site peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADP87509
                                                                                                                                                    reductase and cell cycle progression, to regulate/inhibit cell proliferation and/or to promote apoptosis in a cell in vitro or in vivo. The present sequence represents a thioredoxin family conserved catalytic site peptide, which is given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                 New 4-(1(sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2,5-dienone derivatives are thioredoxin reductase inhibitors useful for treating e.g.
                                                                                                            Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                           Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-517586/49.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-DEC-2002; 2002WO-GB005842
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004056361-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
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                                                                                                                                        invention
                                                                                                                                                                                                                                                                                                                                                       colon or renal cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CANC-) CANCER RES TECHNOLOGY LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-DEC-2002; 2002WO-GB005842
                                                 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WCGPCK 6
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                             WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         family conserved
                                                       Conservative
                                                                                                                                                                                                                                                                                                                           Page 82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                         Poole TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           catalytic site peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 47; DE
Pred. No. 2e+
); Mismatches
                                                                      Score 47; DB 8;
Pred. No. 2e+06;
                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                         Wells G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2e+06;
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                                                         0;
                                                                                  Length 6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Berry JM;
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                                                       ç,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                         Gaps
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                                                                                                                                                                                 CC activity of a pli protein and effects a change in the level of CC plasminogen activation by a cell. Also described: (1) making (M1) a CC clonal cell line, which involves isolating a cell, then characterising CC the activity of a protein produced by the cell or clonal progeny of the CC cell, where the protein is involved in plasminogen activation; and (2) a CC clonal cell line (II) useful in the identification of composition that CC modulate pll activity, where the clonal cell line is obtained by (M1). CC (I) has cytostatic activity, and can be used in antisense therapy. (I) is useful for modulating the activity of pl1 which involves administering CC (I) to a cell. The cell can be a human cancer cell and a CCL-22 colorectal adenocarcinoma cell. (I) is also useful for reducing the development of CC cancer in a patient e.g., mouse which involves administering (I) to a cancer cell in the patient. (I) is useful for inhibiting the growth of CC tumours or inhibiting tumour cell invasion in a patient, which involves administering (I) to a cancer cell in the patient. (II) is useful for inhibiting the composition that modulates pl1 activity which involves cadministering (I) to a cancer cell in the patient. (II) is useful for composition that modulates pl1 activity of a cell of the clonal cell line relative to a call of a cell of the clonal cell line relative to a cancer cell in the patient of (II) obtained by (M1), determining the composition that produces a change in pl1 activity. The composition that produces a change in pl1 activity is a change in the level of plasminogen activation of the present layention.
                             S
                                                                      Matches
                                                                                      Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         activity of pll protein activation by a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Composition useful for inhibiting growth of tumor in patient, modu activity of pl1 protein and effects change in level of plasminogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADP67423 standard; peptide; 6
                                                                                                                                        Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; SEQ ID NO 4; 140pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-487995/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Waisman D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-DEC-2002; 2002US-0433140P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-DEC-2003; 2003WO-US040029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004054517-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADP67423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (WAIS/) WAISMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pll activity modulator; plasminogen activation; cytostatic;
sense therapy; cancer; tumour; tumour growth inhibition; thioredoxin;
                                                                                                                                                                          present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIOMICS LLC WAISMAN D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disulfide isomerase.
                                                                    6
                                                                                   Similarity
                               WCGPCK 6
WCGPCK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and protein disulfide
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        describes a composition (I)
                                                                    0,
                                                                                   Score 47; DB 8;
Pred. No. 2e+06;
                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                isomerase common sequence
                                                                    0
                                                                                                 Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         which modulates
the level of
                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID
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RESULT 7
AD097008
ID AD099
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AC AD09
XX AD09
XX AD09
XX Pilo
CO Synt
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XX Pilo
XX Pilo
XX Pilo
XX Pilo
XX 22-1
XX 28-1
PF 12-1
XX XX Pilo
PR 13-1
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                by a cell. The composition is useful for reducing pl1 protein activity in cancer cells. They are also useful for inhibiting tumour growth or metastasis or treating cancer. The present sequence represents the amino acid sequence of the thioredoxin and protein disulphide isomerase common sequence involved with cleavage, formation and reshuffling of disulphide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New composition which modulates the activity of a pl1 protein and a change in the level of plasminogen activation by a cell useful treating cancer or reducing pl1 protein activity in cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-NOV-2001; 2001US-0333866P
26-NOV-2002; 2002US-00304287
13-DEC-2002; 2002US-0433140P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pl1 protein; plasminogen activation; tumour growth; metastasis; treating.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADQ97008 standard;
                                                                                                                                                                                                                                                                                                                                            AAW45003 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a composition which modulates the activity of a pl1 protein and effects a change in the level of plasminogen activation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; SEQ ID NO 4; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-579971/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Waisman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-DEC-2003; 2003US-00735577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-JUL-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADQ97008;
                                                                                Immunomodulator; immunosuppressant; immunostimulator; treatment;
transplant rejection; autoimmune disease; cancer; infection.
                                                                                                                                                                                                                              27-APR-1998
                                                                                                                                                                                                                                                                                       AAW45003;
                                                                                                                                                                   Immunomodulatory peptide D7233.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (WAIS/) WAISMAN D M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry
                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                         peptide; 12
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Pred. No. 2e+
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 8;
2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 6;
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RESULT 9
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   myasthenia gravis, encephalomyelitis, phemphigus vulgaris, v
foliaceus, Senear-Usher syndrome or Brazilian phemphigus. An
immunostimulator can be used to treat conditions such as can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY09429;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sjogren's syndrome, scleroderma, mixed connective tissue disease, type dermatomyositis, polymyositis, Reiter's syndrome, Behcet's disease, type I diabetes, Hashimoto's thyroiditis, Graves' disease, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 12; Page 155; 183pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding cysteine- or methionine-containing peptide(s) which have immuno:stimulatory or immunosuppressive activity - can be to treat, e.g. cancers, infection, auto:immune disease or transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-526397/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-APR-1996;
23-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-APR-1997;
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                      WPI; 1999-287953/24
                                                                   Bergstrand H,
                                                                                                                                                                                                                  06-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY09429 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rejection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9739023-A1
                                                                                                                                                                                                                                                                                                               WO9919347-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunoactive peptide SEQ ID NO:54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ASTR ) ASTRA
                                                                                                                                                                     10-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                           mmune
                                                                                                                                                                                                                                                                                                                                                                                                                                    mmunoactive;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present
                                                                                                                                                                                                                                                                                                                                                                                                           response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6; Conserv
                                                                                                                     ASTRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  opressant can be used to treat transplant rejection or au e.g. rheumatoid arthritis, systemic lupus erythematosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; ilarity 100.0%; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide is an immunosuppressant or immunostimulator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ΑΒ.
                                                                                                                                                                                                                                                                                                                                                                                                           immunomodulation; immunosuppression;
se; immunoreactive; autoimmune disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96SE-00001422.
96SE-00003469.
                                                                        Eriksson
                                                                                                                                                                     97US-00949024
                                                                                                                                                                                                                  98WO-SE001801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eriksson
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                                                                        Lindvall M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 47;
Pred. No.
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                                                                        Saernstrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2;
                                                                                                                                                                                                                                                                                                                                                                                                                disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    immunostimulation;
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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coding sequence encoding an immunoreactive peptide and further encoding a protein targeting sequence. The nucleic acid is administered to a patient so that its expression product, an immunoactive peptide, modulates an immune response in a patient. The nucleic acid can also be used to treat cancer, either after surgery to remove a portion of the cancer or after ionizing radiation. A cytokine is also administered in conjunction with the nucleic acid. Cells containing the nucleic acid molecule can also be used for treatment. The immunoactive peptide is immunosuppressive and can be used in patients with autoimmune disease. The present sequence represents a specifically claimed immunoactive peptide from the present
The invention relates to a novel human thioredoxin polypeptide, 22105, fragments of it and allelic variants. 22105 is useful for identifying a compound which modulates the activity or expression of the polypeptide and nucleic acid, where the binds to the polypeptide, and the binding is detected by direct detection of a test compound/polypeptide binding, by
                                                                                                                                          Novel thioredoxin polypeptides and polynucleotides for treating disorder associated with aberrant cellular proliferation or differentiation, e.g., cancer and to identify modulators for the
                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cellular proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU08670;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic genes encoding immunoreactive peptides containing cysteine
                                                                                             Disclosure;
                                                                                                                                                                                                        WPI; 2001-589945/66
                                                                                                                                                                                                                                       Curtis RAJ
                                                                                                                                                                                                                                                                                                   07-MAR-2000;
                                                                                                                                                                                                                                                                                                                                  05-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                WO200166756-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ischaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chronic myelogenous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thioredoxin domain markov model sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   methionine
                                                                                                                                                                                                                                                                     (MILL-) MILLENIUM PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WCGPCK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sarcoma; tumour; haemopoetic neoplastic disorder; reukaemia;
elogenous leukaemia; cardiovascular disorder; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA;
                                                                                             Fig 3A; 126pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        describes nucleic acid molecules comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypertension.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            differentiative disorder; cancer; poetic neoplastic disorder; leuka
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for therapeutic
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disorder (e.g. cancers, carcinomas, sarcomas and tumours), haemopoetic neoplastic disorder (e.g. leukaemia and chronic myelogenous leukaemia) or cardiovascular disorder (e.g. atherosclerosis, ischaemia, thrombosis, hypertension), characterised by aberrant activity or expression of a 22105 nucleic acid or polypeptide in a subject, where the agent is preferably a peptide, a phosphopeptide, a small molecule, an antibody or their combination, or an antisense, a ribozyme, a triple helix molecule, a 22105 nucleic acid or their combination (many examples of each disorder are given in the specification). The present sequence represents a thioredoxin domain sequence derived from a hidden markov model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  using a competition binding assay or an assay for 22105-mediated signal transduction, and the activity of the polypeptide is a redox activity or the ability to modulate protein processing, protein folding and protein secretion. Anti-22105 antibody is useful for detecting the presence of 22105 in a sample. A modulator of 22105 is useful for treating or preventing a disorder such as cellular proliferative or differentiative
          A 14 kDa molecule designated various names (Eosinophil Cytotoxicity-Enhancing Factor (ECEF); thioredoxin and ADF) and encoding a polypeptide of 104 amino acids has been found to support the growth of an EBV transformed B lymphocyte cell line in the manner of IL-1; to induce the expression of IL-2 receptors in a large granular lymphocyte cell line from an ATL patient in the manner of IL-1 and to mediate the growth inhibitory properties of interferon gamma. Recombinant ECEF (FECEF) 80 is
                                                                                                                                                                                                                                                                                                                                                            06-APR-1992;
06-JUL-1992;
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04-MAY-1994
                                                                                                                                                   Example 3;
                                                                                                                                                                                 Shortened Eosinophil Cytotoxicity-Enhancing Eosinophil-stimulating activity, for use in
                                                                                                                                                                                                                                                          WPI; 1993-336837/42.
                                                                                                                                                                                                                                                                                          Silberstein DS,
                                                                                                                                                                                                                                                                                                                                                                                                              06-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eosinophil cytotoxicity enhancing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR42822 standard; protein; 80 AA
                                                                                                                                                                                                                                                                                                                          (BGHM ) BRIGHAM & WOMENS HOSPITAL
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                                                                                                                                                   Page 67; 96pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
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92US-00906842
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                                                                                                                                                   English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  factor; ECEF; stimulation; diagnosis;
                                                                                                                                                                                   Factor proteins - having diagnosis and therapy.
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version of the 104 amino acid polypeptide

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                                     A 14 kDa molecule designated various names (Bosinophil Cytotoxicity-Enhancing Factor (ECEF); thioredoxin and ADF) and encoding a polypeptide of 104 amino acids has been found to support the growth of an EBV transformed B lymphccyte cell line in the manner of IL-1; to induce the expression of IL-2 receptors in a large granular lymphccyte cell line from an ATL patient in the manner of IL-1 and to mediate the growth inhibitory properties of interferon gamma. Recombinant ECEF (RECEF) 84 in a truncated version of the 104 amino acid polypeptide comprising the first 84 amino acids of the molecule. Some shortened ECEF's are useful because they retain eosinophil strimulating activity but lack proinflammatory dithiol reductase activity. In particular they can be used for killing tumour cells. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               first 80 amino acids of the molecule. Some shortened ECEF's are useful because they retain eosinophil stimulating activity but lack proinflammatory dithiol reductase activity. In particular they can be used for killing tumour cells. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                           Shortened Eosinophil Cytotoxicity-Enhancing Factor proteins - having Eosinophil-stimulating activity, for use in diagnosis and therapy.
                                                                                                                                                                                                                                                                                                                                                                                                         06-APR-1992;
06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eosinophil cytotoxicity enhancing factor; ECEF; stimulation; diagnosis; therapy; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
04-MAY-1994
                                                                                                                                                                                                                                                                                                                      WPI; 1993-336837/42.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    field.)
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6; Conserv
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92US-00906842.
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Pred. No.
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RESULT 13

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                                                                                                                                                                                for therapeutic intervention in treating a disease associated with altered mitochondrial function. Specifically, it refers to a method for identifying proteins of the human heart mitochondrial proteome that are useful for drug screening assays, as well as therapeutic targets. The present invention describes a method for identifying such proteins that can be used in the treatment of various diseases associated with altered mitochondrial function including diabetes mellitus, Huntington's disease, osteoarthritis, Leber's hereditary optic neuropathy (LHON), mitochondrial encephalopathy lactic acidosis and stroke (MELAS) mycolonic epilepsy
                                                                   Sequence
                                                                                                 ragged red fibre syndrome (MERRF) or cancer. Accordingly, these compositions have neuroprotective, nootropic, antidiabetic, anticonvulsant, antiarthritic, osteopathic, ophthalmological and cytostatic activities. This polypeptide sequence is a human hear mitochondrial protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                             comprises detecting a with the disease.
                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying a mitochondrial target for drug screening assays and for treating diseases associated with altered mitochondrial function, comprises detecting a modified polypeptide in a sample and correlation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-APR-2002; 2002US-0372843P.
17-JUN-2002; 2002US-0389987P.
20-SEP-2002; 2002US-0412418P.
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osteopathic; ophthalmological; cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mitochondrial encephalopathy lactic acidosis and stroke; MELAS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Huntington's disease; osteoarthritis; Leber's hereditary optic neuropathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mitochondrial; human; screening assay; diabetes mellitus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human heat mitochondrial protein as a therapeutic target SeqID312.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                          This invention relates to novel mitochondrial targets that can be used
                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 312; 180pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-845369/78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ghosh SS,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          myoclonic epilepsy ragged red fibre syndrome; MERRF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ68506 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BUCK-)
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Local Similarity
les 6; Conserv
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BUCK INST AGE RES.
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                 100.0%;
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Pred. No.
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                                 Length 84
                                                                                                                    is a human heart
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                                                                                                                                                                                                                                                                                                                                                                                                                              correlating
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Mismatches

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Indels

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Query Match

100.0%;

Score 47;

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                                                     MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to ABP60964 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                       The present invention describes a method (M1) for producing an oil body associated with a recombinant multimeric protein complex (MC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and (P1, P2), where P3 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associated with a recombinant P1 M1 is useful for producing an oil body associated with a recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e.g. redox proteins and polypeptides capable of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oil body; ophthalmological; antidiabetic; cytostatic; antipsoria vasotropic; vulnerary; antibacterial; immunosuppressive; antiulc food product; milk; wheat; oxidative stress; cataract; diabetes;
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04-DEC-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 81; Page 243; 362pp; English
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Val G, Zaplachinski S,
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2001US-0302885P.
2001US-00006038.
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Moloney M;
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sepsis;

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RESULT 15
AAB49948
Search completed: February 23, 2006, 23:11:52
Job time : 187 secs
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                                                                                                      Query Match 100.0%; 9
Best Local Similarity 100.0%; 1
Matches 6; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             17-MAY-1999;
18-MAY-1999;
02-JUN-1999;
                                                                                                                                                                                 The present invention provides variants of the Aspergillus oryzae protein disulfide isomerase enzyme. These are capable of reducing disulphide bonds in proteins, which may be the cause of allergies in humans. The proteins can be used to reduce the allergenicity of foods, to treat scleroproteins, in detergents, in food additives and in cosmetics
                                                                                                                                                                                                                                                                                           Protein disulfide isomerase variant having increased reducing properties and decreased redox potential than native proteins, used to reduce allergenicity of allergic proteins in feed, food or cosmetic products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disulphide bond; allergy; protein disulfide isomerase variant; detergent; food additive; cosmetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB49948 standard; protein; 95 AA.
                                                                                                                                                            Sequence 95 AA;
                                                                                                                                                                                                                                                                 Claim 2; Page 76-77; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                WPI; 2001-070776/08.
N-PSDB; AAC89030.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Aspergillus oryzae.
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99DK-00000689.
99US-0137068P.
                                                                                                       Score 47; DB 4; Length 95; Pred. No. 13; Indels
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 length: 0
length: 2000000000
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Match
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14.803 Million cell updates/sec
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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       thioredoxin - Myco
probable thioredox
thioredoxin TC0826
thioredoxin II - y
thioredoxin II - y
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4 5	44	43	42	41	40	39	38	37	36	35	34	33,	32	31	30
47	47	47	47	47	47	47	47	47	47	47	47	47	47	47	47
100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
117	116	115	113	110	110	110	109	109	109	109	108	108	108	107	107
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E70107	B70851	T29044	857775	E81850	C81090	A87688	AF0922	S27053	TXEC	S35497	D82338	AD0471	G82991	E64047	AH3504
thioredoxin (trxA)	probable trxC prot	hypothetical prote	thioredoxin h, cyt	thioredoxin I NMA1	thioredoxin NMB136	thioredoxin (impor	thioredoxin (impor	thioredoxin - Emer	thioredoxin [valid	thioredoxin - Salm	thioredoxin VC0306	thioredoxin 1 (imp	thioredoxin PA5240	thioredoxin - Haem	thioredoxin C-1 [i

ALIGNMENTS

RESULT 1
E844409
thioredoxin [imported] - Halobacterium sp. NRC-1

C;Species: Halobacterium sp. NRC-1
C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 05-Oct-2004
C;Accession: E84409
R,Ng, W.V.; Kennedy, S.P.; Mahairas, G.G.; Berquist, B.; Pan, M.; Shukla, H.D.; Lasky, S; Leithauser, B.; Keller, K.; Cruz, R.; Danson, M.J.; Hough, D.W.; Maddocks, D.G.; Jablo Jung, K.H.; Alam, M.; Freitas, T.
Proc. Natl. Acad. Sci. U.S.A. 97, 12176-12181, 2000
A;Authors: Hou, S.; Daniels, C.J.; Dennis, P.P.; Omer, A.D.; Ebhardt, H.; Lowe, T.M.; Li A;Title: Genome sequence of Halobacterium species NRC-1.
A;Reference number: A84160; MUID:20504483; PMID:11016950
A;Accession: E84409

A;Status: preliminary A;Molecule type: DNA A;Residues: 1-89 <5TO> A;Cross-references: UNIPROT:Q9HMD0; UNIPARC:UPI0000063B9A; GB:AE004437; NID:g10581982; P C;Genetics:

					,	•		,				
C;Genetics: A;Genetic code: SGC3 C;Superfamily: thioredoxin; thioredoxin homology	A;RESIQUES: 1-102 <11GK> A;CCG08-references: UNIPROT:P47370; UNIPARC:UPI0000136E89; GB:U39691; GB:L43967; NID:g10 A;Experimental source: strain G-37	A;Status: nucleic acid sequence not shown; translation not shown A;Molecule type: DNA A;Molecule type: DNA	A,Title: The minimal gene complement of Mycoplasma genitalium. A,Reference number: A64200; MUID:96026346; PMID:7569993 A,Accession: G64213	, C.A.; Venter, J.C. Science 270, 397-403, 1995	R;Fraser, C.M.; Gocayne, J.D.; White, O.; Adams, M.D.; Clayton, R.A.; Fleischmann, R.D.; M.; Fuhrmann, J.; Nguyen, D.; Utterback, T.R.; Saudek, D.M.; Phillips, C.A.; Merrick, J.	C;Date: 17-Nov-1955 #sequence_revision 16-Aug-1996 #text_change 05-Oct-2004 C;Accession: G64213	thoredoxin - Mycoplasma genitalium	RESULT 2	Db 12 WCGPCK 17	Qy 1 WCGPCK 6	Query Match 100.0%; Score 47; DB 2; Length 89; Best Local Similarity 100.0%; Pred. No. 0.93; Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	A;Gene: trxA2 C;Superfamily: thioredoxin

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A; Molecule type: DNA
A; Residues: 1-102 <HIM>
A; Cross-referen
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A;Note: the nucleotide sequence was
C;Genetics:
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C; Superfamily: thior
F; 9-90/Domain: thior
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A;Title: Complete sequence analysis of the genome of the bacterium Mycoplasma A;Reference number: S73327; MUID:97105885; PMID:8948633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Mycoplasma pneumoniae
A;Vartlety: ATCC 20342
C;Date: 27-Feb-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Accession: S73896
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                                                                                                                                                                                                                                                                                                                                          R;Stephens, R.S.; Kalman, Science 282, 754-759, 1998
                                                                                                                                                                                                                                                                                                                                                                               C;Date: 13-Sep-1998 #sequence_revision 13-Sep-1998 #text_change 05-Oct-2004
C;Accession: B71503
                                                                                                                                                                                                                                                                                                                                                                                                                 probable thioredoxin - Chlamydia trachomatis (serotype D, strain UW3/Cx) C_iSpecies: Chlamydia trachomatis
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                                                                                                                                          Superfamily: thioredoxin; thioredoxin homology
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Best Local
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100.0%; Pred. No.
                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                               S.; Lammel, C.J.; Fan, J.; Marathe, R.; Aravind, L.; Mitchell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47;
                                                                     Pred. No. 1;
; Mismatches
                                                                                                                                                                                                 UNIPARC:UPI0000136E75; GB:AE001324; GB:AE001273; strain UW-3/Cx
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              UNIPARC: UPI0000136E8A; EMBL: AE000056; submitted to the EMBL Data Library, No
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R;Gan, Z.R.
R;Gan, Z.R.
J. Biol. Chem. 266, 1692-1696, 1991
J. Rivitle: Yeast thioredoxin genes.
A;Title: Yeast thioredoxin genes.
A;Reference number: A38669; MUID:91
A:Accession: S15048
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C;Species: Chlamydia muridarum, Chlamydia trachomatis MoPn

C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 05-Oct-2004

C;Accession: C81660

R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hickey, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzberg, Nucleic Acids Res. 28, 1397-1406, 2000

Nucleic Acids Res. 28, 1397-1406, 2000

A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.

A;Reference number: A81500; MUID:20150255; PMID:10684935
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A; Residues: 1-103 < MUL>
A; Residues: 1-103 < MUL>
A; Cross-references: UNIPARC: UPI00004F943;
A; Cross-references: UNIPARC: Holmgren, A;
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A;Experimental source: strain Nigg (
C;Genetics:
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Eur. J. Biochem. 23, 328-335,
A;Title: Yeast thioredoxin. Ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          J. Biol. Chem. 266, 9194-9202, 1991
A; Title: Thioredoxin deficiency in yeast prolongs S phase A; Reference number: A39847; MUID:91225027; PMID:2026619
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C;Species: Saccharomyces cerevisia
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                                                                                                                                                                                                                                                                                                                                                                                      R; Koetter, P.; Rose, M.; Entian, K.D. submitted to the Protein Sequence Database,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Reference number: A39847;
A;Accession: A39847
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A;Cross-references: UNIPROT:P22217; UNIPARC:UPI000004F943;
A;Gene: SGD:TRX1; TR-II; MIPS:YLR043c
A;Cross-references: SGD:S0004033; MIPS:YLR043c
                                                                                                                   A; Note: experimental_source
                                                                                                                                                                    A; Cross-references: UNIPARC: UPI000004F943;
                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-103 < KOE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Accession: S15360
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100.0%; Pred. No.
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                                                                                                                                                                    EMBL: Z73215; NID: g1360372; PIDN: CAA97572.1;
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C;Superfamily: thioredoxin; thioredo
C;Keywords: redox-active disulfide
F;2-103/Product: thioredoxin II #st:
F;9-90/Domain: thioredoxin homology
F;30-33/Disulfide bonds: redox-activ
                                                                         A;Cross-references: UNIPROT:P08628; UNIPARC:UPI0000136E96 C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                        J. Biol. Chem. 263, 9589-9597, 1988
J. Biol. Chem. 263, 9589-9597, 1988
A;Title: Amino acid sequence of thioredoxin isolated from A;
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R;Johnson, R.S.; Mathews, W.R.; Biemann, K.; Hopper,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Superfamily: thioredoxin; thioredoxin homology
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A;Accession: T46567
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;Cross-references: UNIPROT:014463; UNIPARC:UPI000016205F; EMBL:Z99532; PIDN:CAB16724.1; Experimental source: strain 972h-; cosmid c7D4; Experimental source: strain 972h-; cosmid c7D4; Lenaers G.; Perret, E.; Bonin, O.; Picaerd, A.; Caput, D. ubmitted to the EMBL Data Library, October 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reference number: Z21826
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Species: Oryctolagus cuniculus (domestic rabbit)
Date: 30-Jun-1989 #sequence_revision 26-May-1994 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Experimental source: strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ubmitted to the EMBL Data Library, October 1997; Description: TRX2, a fission yeast stress protein.; Reference number: Z23075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;Residues: 1-103 <GEN>
                                                                                                                                                          Residues: 1-104 <JOH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gene: trx2; SPAC7D4.07c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Accession: T39085; T46567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Species: Schizosaccharomyces pombe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Alternate names: thioredoxine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Status: preliminary; translated from GB/EMBL/DDB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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ce: strain 972(-)
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homology <THR>
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Pred. No. 1;
0; Mismatches
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Pred. No.
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Nature 390, 249-256, 1997

A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Galler, A;Authors: Foulger, D.; Fritz, C.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F. iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F. Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois, A;Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueel Y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon, A; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Seror akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchlyama, T.; Winters, P.; Wipat, A.; Yamanoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K.; A;Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.
                          R;Gan,
                                                                                             thioredoxin I - yeast (Saccharomyces cerevisiae) N;Alternate names: protein G7746; protein YGR209c; C;Species: Saccharomyces cerevisiae
                                                                                                                                                                                                RESULT 10
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R;Kunst, F; Ogasawara, N; Moszer, I; Albertini, A.M.; Alloni, G; Azevedo, V; Berter
C.; Bron, S; Brouillet, S; Bruschi, C.V.; Caldwell, B; Capuano, V; Carter, N.M.; Cho
A; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J; Fabret, C.; Ferrari, E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Chen,
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                                                                                                                                                                           TXBY1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Keywords: redox-active disulfide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Map position:
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A;Residues: 1-104 <KUN>
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                                                                      Date: 31-Dec-1991
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Best Local
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;Date: 31-Jan-1992 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
                                                                                                                                                                                                                                                                                                                                                                            Matches
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Ehrlich,
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piol. 135, 2931-2940, 19
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266, 1692-1696, 1991
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53932; S61947; S64531; S63858; A38669
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A; Molecule type:
A; Residues: 1-104
A; Cross
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A; Molecule type: protein

A; Molecule type: protein

A; Residues: 2;27-43;98-104 <HAL>

A; Residues: 2;27-43;98-104 <HAL>

A; Residues: 2;27-43;98-104 <HAL>

A; Cross-references: UNIPARC: UPI000011EC95; UNIPARC: UPI0000171E1F; UNIPARC: UPI0000171E20

A; Cross-references: Tommary and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with that from page and from Fig. 5 is inconsistent with the figure of the fig. 5 is inconsistent with the figure of the
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A;Residues: 1-104 <GUE>
A;Residues: 1-104 <GUE>
A;Residues: 10104 <GUE>
A;Cross-references: UNIPARC:UPI000004F91C; EMBL:Z49133; NID:g790489; PIDN:CAA89002.1;
A;Experimental source: strain S288C
A;Experimental source: strain S288C
B;Song, J.M.; Cheung, E.; Rabinowitz, J.C.
submitted to the EMBL Data Library, November 1995
submitted to the EMBL Data Library, November 1995
A;Description: Analysis of the 15.6-kb fragment encompassing the ADE3 gene.
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A;Accession: B39847
A;Molecula - ---
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R;Guerreiro, P.; Barreiros, T.; Soares, H.; Cyrne, L.; Maia e Silva, A.; Rodrigues-Pous: R;Guerreiro, P.; Barreiros, T.; Soares, H.; Cyrne, L.; Maia e Silva, A.; Rodrigues-Pous: Yeast 12, 273-280, 1996
Yeast 12, 273-280, 1996
A;Title: Sequencing of a 17.6 kb segment on the right arm of yeast chromosome VII reveal terial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia coll phosterial electron-transferring flavoproteins (beta-chain) and of the Escherichia electron-transferring flavoproteins (beta-chain) and the electron-transferring f
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R;Hall, D.E.; Baldesten, A.; Holmgren, A.;
Eur. J. Biochem. 23, 328-335, 1971
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J. Biol. Chem. 266, 9194-9202, 1991
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A;Reference number: A38669; MUID:
A;Accession: S15049
A;Molecule type: DNA
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C; Superfamily: t
C; Keywords: redc
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A; Accession: S64531
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A;Residues: 1-104 <S
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A; Accession: S53932
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Residues: 1-104 <MUL>
Best Local Similarity
                                                              Query Match
                                                                                                                                                                                                                                            9-91/Domain: thioredoxin homology <THR>
                                                                                                                                                                                                                                                                                             2-104/Product: thioredoxin I #status experimental
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Score 47;
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                                                 DB 1;
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                                                       Length 104;
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C;Accession: B59394; A59357
R;Reynolds, C.M.; Meyer, J.; Poole, L.B.
Rinchemistry 41, 1990-2001, 2002
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A;Title: Exploring the Mycoplasma capricolum genome: a minimal A;Reference number: S77739; MUID:96059641; PMID:7476192

A;Accession: S77780
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Experimental source: ATCC 2734
A;Note: the nucleotide sequence
C;Genetics:
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                                                                                     A;Cross-references: UNIPARC:UPI0000174DEF
R;Hammel, K.E.; Cornwell, K.L.; Buchanan, B.B.
Proc. Natl. Acad. Sci. USA 80, 3681-3685, 1983
A;Title: Ferredoxin/flavoprotein-linked pathway for the re
A;Reference number: A59432
A;Contents: annotation; purification and characterization
                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPARC:UPI0000174DEF R;Reynolds, C.M.; Poole, L.B.; Hammel, K.E.; Wada, K.; Buchanan, submitted to the Protein Sequence Database, September 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-104 <BOR>
                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: protein A; Residues: 1-104 < RE1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Clostridium pasteurianum
C;Date: 02-Apr-2002 #sequence_revision 02-Apr-2002 #text_change 05-Oct-2004
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                                                                                                                                                                                                                    A; Molecule type: protein A; Residues: 1-104 < RE2>
                                                                                                                                                                                                                                                               A;Status: preliminary
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                                                                                                                                                                                                                                                                                                         A; Reference number: A59394
                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Accession: B59394
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thioredoxin - Clostridium pasteurianum
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mes: protein MC064
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Pred.
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                                                                            RESULT 15
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                                        thioredoxin - chicken
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ancet 357, 1225-1240, 20
;Species: Gallus gallus (chicken)
;Date: 31-Mar-1989 #sequence_revi
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Best Local (
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Reference number: A
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Experimental source: strain C-125
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Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 05-Oct-2004
Accession: B84037
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                                                                                                                                         28
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source: strain N315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       me sequencing of meticillin-resistant Stapylococcus aureus. A89758; MUID:21311952; PMID:11418146
#sequence_revision 26-May-1994 #text_change 05-Oct-2004
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4317-4331, 2
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                                                                                                                                                                                                                                  100.0%; Score 47; DB 2; 100.0%; Pred. No. 1;
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K.;
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u. Blol. Chem. 263, 9607-9611, 1988
A;Title: Isolation of a chicken thioredoxin cDNA clone: thioredoxin mRNA is differential
A;Reference number: A30006; MUID:88257080; PMID:2838473
A;Accession: A30006
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                             S
                                                                                                                                   F;32-35/Disulfide bonds:
                                                                                                                                                                                                                                  A; Molecule type: mRNA
                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                  Residues: 1-105 <JON>
                                                                  Matches
                                                                                                                                                     9-92/Domain:
                                                                                                                                                                    Keywords: redox-active disulfide
                                                                                                                                                                                   Superfamily: thioredoxin;
                                                                                                                                                                                                  Cross-references: UNIPROT: P08629; UNIPARC: UPI00001713BC; GB: J03882; NID: g212765; PIDN:
 31
                                                                6,
WCGPCK 36
                               WCGPCK 6
                                                                                                                                                 thioredoxin homology <THR>
                                                                Conservative
                                                                                                                                 redox-active #status predicted
                                                                                  100.0%;
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Pred. No.
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                                                                                                1; Length 105
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Search completed: February 23, Job time : 40 secs

2006, 23:16:36

Mis Page Blooms (1870).

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Result
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Minimum DB seq length: 0
Maximum DB seq length: 200000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                             Database
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence:
                                                                                                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                         Score
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Match Length DB
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1: uniprot_sprot:*
2: uniprot_trembl:*
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  GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd.
                                                                                                                                                                                                                                     Q4T7B7 TETNG
060744 HUMAN
Q5T936 HUMAN
Q5V0J2 HALMA
Q5V0J2 HALMA
Q9HWD0 HALSA
THIO THIRO
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THIO CHLMU
THIO CHLMU
THIO CHLMU
THIO MYCGE
THIO MYCPN
THIO SCHPO
TRX1 YEAST
Q51/18 MESFL
Q61/18 MESFL
Q60/100 MYCMS
                                                                                                                                                    Q4XKV3_PLACH
Q4ES18_PLABE
Q4ES18_PLABE
Q5LB12_BACFN
Q64RG1_BACFN
Q8AB91_BACTN
THIO_MYCGA
Q9AB91_CHAYCGA
THIO1_CHLTE
Q6K1B7_MYCMO
THIO BĀCSU
TRX2 YEAST
Q75CQO ASHGO
Q6FND5 CANGA
Q66J80 DEBHA
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Q64rg1
Q8ab91
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Q6mug0
P14949
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•	Query Matc Best Local Matches	G1212 H , 3 2 H.	NUCLEOTIDE Jaillon O. Mauceli E. Micaud S. Nicaud S. Dasilva C. Biemont C. Cruaud C. Cruaud C. Cruaud C. Lindblad-Tr	13-SEP-200 13-SEP-200 13-SEP-200 13-SEP-200 Chromosome (Fragment) ORFNames=G Tetracdon Eukaryota; Actinopter Acanthomor Tetradonto ORGEI_TaxID	JLT 1 787 TETNG Q4T787	\$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$ \$
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	<pre>k; Score 47; DB 2; Length k; Pred. No. 2; 0; Mismatches 0; Indel</pre>	ute Centre for Genome Rese EMBL/GenBank/DDBJ databas hown here is derived from genome shotgun (WGS) entl 5.1; -; Genomic_DNA. 64705ABEA4A5620E CRC64;	st F., Petit J.L., Stange Sther C. Ozouf-Costaz C., S., Luffalla G., Dossat S., Luffalla G., Dossat Pelli V., Katinka M., Vaclica L., Poulain J., De ler P., Coutanceau J.P., Le C., McKernan K.J., McE R., Zody M.C., Mesirov J. Nusbaum C., Kahn D., Robstier F., Saurin W., Scarissenbach J., Roest Crollicator fish Tetraodon nicaryotype.";	Created) Last sequence update) Last sequence update) Last annotation update) F8161, whole genome shotg en puffer). en puffer). graniata; Vertebrata; freleostei; Euteleostei; gii; Percomorpha; Tetraod dae; Tetraodon.	ALIGNMENTS PRT; 80 AA.	Q7PAWB_FUSNV Q5FLW1_LACAC Q8RH23_FUSNN Q74KU7_LACJO THIO_BOVIN THIO_CALJA THIO_CHICK THIO_HORSE THIO_HUMAN THIO_MOLSE THIO_MOLSE THIO_MOLSE THIO_POHHA THIO_POHHA THIO_POHHA THIO_POHHA THIO_POHHA THIO_RABBIT
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Matches 6
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01-FEB-2005
01-FEB-2005
                                      Submitted (MAY-2005) to the EMBL/GenBank/DDBJ databas EMBL; AL158158; CAI14067.1; -; Genomic_DNA.

SMR; Q57936; 1-85.

GO; GO:0005489; F:electron transporter activity; IEA.

GO; GO:0006118; P:electron transport; IEA.

InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredoxin-1ike.

Pfam; PF00085; Thioredoxin; 1.
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Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
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01-MAR-2004 (TrEMBLrel. 26, Las
Thioredoxin delta 3 (Fragment).
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6 HUMAN
Q5T936 HUMAN PRELIMINARY;
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GO; GO:0005489; F:electron transporter activity; GO; GO:0005189; P:electron transport; IEA.

InterPro; IPR006662; Thiored.

InterPro; IPR012336; Thioredox dom2.

InterPro; IPR012336; Thioredoxin_like.

InterPro; IPR012335; Thioredoxin_fold.
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TISSUE=Liver;
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01-AUG-1998
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Eukaryota; Metazoa;
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PRINTS; PR00421; THIOREDOXIN
PROSITE; PS00194; THIOREDOXIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wang Y., Wang Y.G., Zhang Y., Yuan Y., Nan alternative splice variant of human Chin. Sci. Bull. 43:292-295(1998).
EMBL; AF065241; AAC17430.1; -; mRNA.
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                        PRINTS;
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                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=TXN; ORFNames=RP11-427L11.1-002;
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RESULT 5

Q9HMD0 HALSA
ID Q9HMD0 H
AC Q9HMD0 T
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OR Thioredo
GN Name=trx
OS Halobact
OC Archaea;
OC Halobact
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Q9HMD0;
01-MAR-2001 (
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GMBL; AY596297; AAV46961.1; -; Genomic GO; GO:0005489; F:electron transporter GO; GO:0006118; P:electron transport; I InterPro; IPR006662; Thiored.

InterPro; IPR005746; Thioredoxin.

InterPro; IPR006663; Thioredox dom2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
STRAIN=NRC-1 / ATCC 700922 / JCM 11081;
MEDLINE=20504483; PubMed=11016950; DOI=10.1073
                                                           Halobacterium salinarium (Halobacterium halobium).
Archaea; Euryarchaeota; Halobacteria; Halobacteriales;
Halobacteriaceae; Halobacterium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Baliga N.S., Bonneau R., Facciotti M.T., Pan N
Deutsch E.W., Shannon P., Chiu Y., Weng R.S.,
Date S.V., Marcotte E., Hood L., Ng W.V.,
"Genome sequence of Haloarcula marismortui: a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Name=trxA3; OrderedLocusNames=rrnAC2110;
Haloarcula marismortui (Halobacterium ma
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PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=ATCC 43049;
PubMed=15520287; DOI=10.1101/gr.2700304;
                                                                                                          Thioredoxin
                                                                                                                      01-MAR-2004
                                                                                                                                01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                TIGRFAMs; TIGR01068; thioredoxin;
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                                                                                               Name=trxA2; OrderedLocusNames=VNG2600G;
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(TrEMBLrel. 16,
(TrEMBLrel. 26,
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  DOI=10.1073/pnas.190337797;
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P96132;
15-JUL-1998
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                                                                                                                                                                                                                                                                                                     This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Thiocapsa roseopersicina. Bacteria; Proteobacteria; Chromatiaceae; Thiocapsa.
                                                                                                                                     EMBL; U75512; AAB36882.1;
HSSP; P00274; 2TRX.
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Submitted (AUG-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=M1;
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TIGRFAMS; TIGR01068; thioredoxin;
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InterPro; IPR006662; Thiored.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR006663; Thioredox_dom2.
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EMBL; AE005133; AAG20641.1; -; Genomic_DNA.
            InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
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                                                                                                                                                                                                                                                                                                                                                                                                   similarity)
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Pred. No. 2.2;
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Quail M.A., Ormond D., Doorett J
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PROSITE; PS00194; THIOREDOXIN; 1.

Electron transport; Redox-active center;

DISULFID 33 36 Redox-active

NON_TER 91 91
                                                           Q4Z518 PLABE
Q4Z518;
13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
Thioredoxin, putative (Fragment)
ORFNames=PB000593.00.0;
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NON TER 1
SEQUENCE 95 AA; 1
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                                                                                                                                                                                                                                                                                                                                                                                                    transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                           Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J., Blidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos Janes C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.; "A comprehensive survey of the Plasmodium life cycle by genomic,"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
Hall N., Karras M.,
Berriman M., Florens
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NCBI_TaxID=5825;
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EMBL/GenBank/DDBJ whole genome shotgun (WGS)
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Pred. No. 2.3;
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Pred. No.
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n A., Christophides
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RESULT
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Best Local Similarity
Matches 6; Conserv
                         InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
TIGRPAMS; TIGR01068; thioredoxin; 1.
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Eukaryota; Alveol
NCBI_TaxID=5821;
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                                                                                                                                                                               Cerdeno-Tarraga A.-M., Patrick S., Crossman L.C., Blakely G., Abratt V., Lennard N., Poxton I., Duerden B., Harris B., Quai Barron A., Clark L., Corton C., Doggett J., Holden M.T.G., La Line A., Lord A., Norbertczak H., Ormond D., Price C., Rabbinowitsch E., Woodward J., Barrell B.G., Parkhill J.; "Extensive DNA inversions in the B. fragilis genome control v gene expression.";
                                                                                                                                                                                                                                                                                                                                                                                                                              01-FEB-2005 (TrEMBLrel. 29, Created)
01-FEB-2005 (TrEMBLrel. 29, Last sequence up
01-FEB-2005 (TrEMBLrel. 29, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preliminary data; CAH94443.1; -; Gen EMBL; CAAI01000448; CAH94443.1; -; Gen InterPro; IPR006662; Thioredo. InterPro; IPR006663; Thioredox.dom2. InterPro; IPR012336; Thioredox.in-like. InterPro; IPR012335; Thioredox.in-fold. Pfam; PF00085; Thioredoxin; 1.
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Bacteroidaceae; Bacteroides.
NCBI_TaxID=272559;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transcriptomic, and proteomic analyses."; Science 307:82-86(2005)
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                                                                                                                                                      Science 307:1463-1465(2005).
EMBL; CR626927; CAH08710.1;
                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA]. PubMed=15746427; DOI=10.1126/science.1107008;
                                                                                                                                                                                                                                                                                                                                                                       Bacteroides fragilis (strain ATCC 25285 / NCTC 9343).
Bacteria; Bacteroidetes; Bacteroidetes (class); Bacteroidales;
                                                                                                                                                                                                                                                                                                                                                                                                        OrderedLocusNames=BF3015;
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            TIGRFAMS; TIGR010
PROSITE; PS00194;
                                                                                                                                                                                                                                                                                                                                                                                                                      Putative thioredoxin
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Pred. No. 2.3;
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A., Christophides G
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RESULT 11

QBAB91 BACTN

ID AB51B1 BACTN

AC QBAB91;

DT 01-JUN-2003 (T)

DT 01-WAR-2004 (T)

DT 01-WAR-2004 (T)

DE Thioredoxin C-GN OrderediocusNa

OS Bacteroides th

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Q64RG1_BACFR
ID Q64RG1,
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Matches 6
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Best Local S
Matches 6
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K KUWAhara T., Yamashita A., Hirakawa H., Nakayama H., Toh H., Ok Kuwahara T., Yamashita A., Hirakawa H., Nakayama H., Toh H., Ok Kuwahara S., Hattori M., Hayashi T., Ohnishi Y.;

"Genomic analysis of Bacteroides fragilis reveals extensive DNA interesions regulating cell surface adaptation.";

Proc. Natl. Acad. Sci. U.S.A. 101:14919-14924 (2004).

EMBL; AP006841; BAD49920.1; -; Genomic_DNA.

R GO; GO:0005148; F:electron transporter activity; IEA.

R GO; GO:0006118; P:electron transport; IEA.

R GO; GO:0006118; P:electron transport; IEA.

R InterPro; IPR006662; Thiored.

R InterPro; IPR006663; Thioredoxin.

R InterPro; IPR012336; Thioredoxin.

R InterPro; IPR012335; Thioredoxin.-like.

InterPro; IPR012335; Thioredoxin.-like.

InterPro; IPR012335; Thioredoxin.-like.
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                                                                                                        Bacteroides thetaiotaomicron.
Bacteria; Bacteroidetes; Bact
Bacteroidaceae; Bacteroides.
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Bacteria; Bacteroidetes; Bacteroidetes (class); Bacteroidales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OrderedLocusNames=BF3175;
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Q64RG1;
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     NUCLEOTIDE SEQUENCE.
STRAIN=VPI-5482 / ATCC 29148;
                                                                                                                                                                                 OrderedLocusNames=BT0219;
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                                                                           NCBI_TaxID=818
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Nguyen D.K., Swartzell S., Madan A., Mahairas
"The complete genome sequence of the avian pat
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GO; GO:0005489; F:electron transporter act: GO; GO:0006118; P:electron transport; IEA.
GO; GO:0006810; P:transport; IEA.
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                                                                                                                                      Microbiology 149:2307-2316(2003).
-!- FUNCTION: Participates in various redox reactions reversible oxidation of its active center dithiol
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                                                                                                                                                                                                                                                                               MEDLINE=21956528; PubMed=11959450;
Skamrov A.V., Feoktistova E.S., Gol'dman M.A.,
"Mycoplasma gallisepticum rpoA gene cluster.";
FEMS Microbiol. Lett. 208:281-285(2002).
                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
STRAIN=A5969Var.B;
                                                                                                                                                                                                                                                                                                                                                                                     Mycoplasma gallisepticum.
Bacteria; Firmicutes; Mollicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                Name=trxA; Synonyms=trx; ORFNames=MGA_0452;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Thioredoxin (Trx)
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28-FEB-2003
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PRINTS; PR00421; THIOREDOXIN.
TIGRFAMs; TIGR01068; thioredoxin;
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InterPro; IPR006662; Thiored.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR00663; Thioredox dom2
Pfam; PF00085; Thioredoxin; 1.
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Science 299:2074-2076(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Xu J., Bjursell M.K., Himrod J., Deng S., Carmichael L.K., Chiang H.C., Hooper L.V., Gordon J.I.;
"A genomic view of the human-Bacteroides thetaiotaomicron symbiosis.";
                                                                                                                                                                                                                               MEDLINE=22830409; PubMed=12949158; DOI=10.1099/mic.0.26427-0;
                                                                                                                                                                                                                                           STRAIN=R (low)
                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=2096;
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                                                                                      SIMILARITY: Belongs to the thioredoxin family SIMILARITY: Contains 1 thioredoxin domain.
                                  European
                                                                                                  similarity).
SIMILARITY: Belongs
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                                                 Swiss-Prot entry is copyright. It is produced through sen the Swiss Institute of Bioinformatics and the EN
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DISULFID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=92038016; PubMed=1718781; DOI=10.1016/0014-5793(91)81106-I; Skamrov A.V., Bibilashvili R.S.; Skamrov B.V., Bibilashvili R.S.; "Mycoplasma gallisepticum strain S6 genome contains three regions "Mycoplasma gallisepticum strain S6 genome contains three regions "Mycoplasma gallisepticum strain S6 genome contains three regions "Mycoplasma" with 16 S rRNA and two regions hybridizing with 23 S and Theorem 18 and 18 an
                      GO; GO:0005489; F:electron transporter GO; GO:0006118; P:electron transport; InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR005746; Thioredoxin. InterPro; IPR012336; Thioredoxin-like.
                                                                                                                                                 EMBL; L35043; AAF36768.1; -; HSSP; P80579; 1NW2.
                                                                                                                                                                                           Skamrov A., Feoktistova E., Submitted (FEB-2003) to the
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Bacteria; Firmicutes; Mol
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01-OCT-2000 (TrEMBLrel.
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InterPro; IPR012335; Thioredoxin fold.
InterPro; IPR011594; Thioredoxin like.
Pfam; PF00085; Thioredoxin; 1.
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EMBL; AE016969; AAP56983.1; ALT_INIT; Genomic_DNA.
HSSP; P80579; 1QUW.
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TIGR01068; thioredoxin; 1.
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EMBL/GenBank/DDBJ databases
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S -> T (in Ref. 1).
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RA Eisen J.A., Nelson K.E., Paulsen I.T., Heidelberg J.F., Wu M.,
RA Dodson R.J., DeBoy R.T., Gwinn M.L., Nelson W.C., Haft D.H.,
RA Hickey E.K., Peterson J.D., Durkin A.S., Kolonay J.F., Yang F.,
RA Hickey E.K., Peterson J.D., Durkin A.S., Kolonay J.F., Yang F.,
RA Holt I.E., Umayam L.A., Mason T.M., Brenner M., Shea T.P.,
RA Radune D., Vamathevan J.J., Khouri H.M., White O., Gruber T.M.,
RA Radune D., Vamathevan J.J., Khouri H.M., White O., Gruber T.M.,
RA Ketchum K.A., Venter J.C., Tettelin H., Bryant D.A., Fraser C.M.;
T'The complete genome sequence of Chlorobium tepidum TLS, a photosynthetic, anaerobic, green-sulfur betterium.';
T'Toc. Natl. Acad. Sci. U.S.A. 99:9509-9514(2002).
T. FUNCTION: Participates in various redox reactions through the Creversible oxidation of its active center dithiol to a disulfide .

C. I. SIMILARITY: Belongs to the thioredoxin family.

C. I. SIMILARITY: Belongs to the thioredoxin family.
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PRINTS; PR00421; THIOREDOXIN.
PRODOM; PD003679; Thioredoxin_like; 1.
TIGRFAMS; TIGR01068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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28-FEB-2003 (Rel. 41, Last sequence up
13-SEP-2005 (Rel. 48, Last annotation
Thioredoxin 1 (Trx-1).
                                              PROSITE; PS00194; THIOREDOXIN; 1.

Complete proteome; Electron transport; Redox-active center; DISULFID 25 28 Redox-active (By similarity).
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; ThioredoxIn.-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
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STRAIN=TLS / ATCC 49652 / DSM 12025;
MEDLINE=22103685; PubMed=12093901; D
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Pfam; PF00085; Thioredoxin; 1.
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                                                                                                TIGRFAMs;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chlorobium
                                                                                                                                                                                                                                                                                                                                                                                      -!- SIMILARITY: Contains 1 thioredoxin domain.
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                                                                                              PR00421; THIOREDOXIN.
s; TIGR01068; thioredoxin; 1.
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                                 101 AA;
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                                                                                                                                                                                                                                                                                                          content
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                                 11247 MW;
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   Score 47;
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                                 BA78E5511900B754 CRC64;
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 DB 1;
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RESULT 15
QGKIET, MYCMO PRELIMINARY;
AC Q6KIET,
DT Q5-UUL-2004 (TrEMBLrel. 27,
DT 05-UUL-2004 (TrEMBLrel. 27,
DE Thioredoxin.

GN Name=trxà, OrderedLocusName
OS MYCOplasma mobile.

QC Bacteria, Firmicutes; Molli
QX NCBI_TAXID=2118;
RN 011
QX YEARIN=163K / ATCC 43663;
RX PubMed=15289470; DOI=10.110
QX STRAIN=163K / ATCC 43663;
RX PubMed=15289470; DOI=10.110
QX STRAIN=163K / ATCC 43663;
RX PubMed=15289470; DCI=10.110
QX STRAIN=163K / ATCC 43663;
RX Complete proteome.
SQ SEQUENCE 101 AA; 11700 M
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Matches 6
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Matches 6; Conserv
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GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "The complete genome and proteome of Mycoplasma mobile."; Genome Res. 14:1447-1461(2004).
EMBL; AE017337; AATZ7629.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jaffe J.D., Stange-Thomann N., Smith C., DeCaprio D., Fisher S., Butler J., Calvo S., Elkins T., FitzGerald M.G., Hafez N., Kodira C.D., Major J., Wang S., Wilkinson J., Nicol R., Nusbaum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mycoplasma mobile.
Bacteria; Firmicutes; Mollicutes; Mycoplasmataceae; Mycoplasma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=163K / ATCC 43663;
PubMed=15289470; DOI=10.1101/gr.2674004;
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Birren B., Berg H.C., Church G.M.,
                                                                                                                                                                                                                                           PROSITE; PS00194; THIOREDOXIN;
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Search completed: February 23, 2006, 23:15:53

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Result
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Maximum Match 100%
Listing first 45 summaries
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   length: 0
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Match Length
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10.554 Million cell updates/sec
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(c) 1993 - 2006 Biocceleration Ltd.
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US-08-08-26-910-4
US-09-601-144-68
US-09-513-999C-8037
US-08-386-729A-10
US-07-921-848-22
US-07-921-848-22
US-08-165-301A-22
US-08-165-301A-28
US-08-165-301A-29
US-09-165-301A-29
US-09-165-301A-
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US-08-627-907A-6
US-07-901-713A-3
US-08-750-142B-54
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Sequence 54, Appl
Patent No. 5210073
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US-09-107-433-5104	US-09-583-110-5222	US-09-272-342B-6	US-09-270-767-47690	US-09-270-767-32473	US-09-741-243C-4	US-08-775-978-3	US-08-775-978-1	US-09-166-966E-8	US-09-328-352-5360	US-09-765-272A-40	US-09-765-271-40	US-09-536-784-40	US-08-961-083-40	US-09-272-342B-5	US-09-949-016-9663	US-09-489-039A-8017	US-U9-543-681A-439/
Sequence 51	Sequence 52	Sequence 6,	Sequence 47	Sequence 32	Sequence 4,	Sequence 3,	Sequence 1,	Sequence 8,	Sequence 50	Sequence 40	Sequence 40	Sequence 40	Sequence 40	Sequence 5,	Sequence 96	Sequence 80	sequence 4.
104, Ap	5222, Ap	6, Appli	47690, A	32473, A	, Appli	, Appli	Appli	, Appli	5360, Ap	0, Appl	0, Appl	•	0, Appl	, Appli	9663, Ap	8017, Ap	4391, Ap

ALIGNMENTS

RESULT 1 US-08-883-804-5

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Patent No. 5985261

GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: Das, Kumuda C.
TITLE OF INVENTION: USE OF THIOREDOXIN-LIKE MOLECULES FOR
TITLE OF INVENTION: INDUCTION OF MMSOD TO TREAT OXIDATIVE DAMAGE
NUMBER OF SEQUENCES: 5
                                                                Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                              CLASSIFICATION: 514
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: CONNEL1, Gary J.
REGISTRATION NUMBER: 28.020
REFERENCE/DOCKET NUMBER: 287;
TELECOMMUNICATION INFORMATION:
TELEPHONE: 303/863-9700
                                                                                                                                                                                                            TELEFAX: 303/863-0223
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 80203
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.
ZIP: 80203
                                                                                                                                                                                             TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
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)GY: linear
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WCGPCK
                                  WCGPCK 6
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                                                                  100.0%; ilarity 100.0%; Conservative 0
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                                                                                      Score 47; DB 1;
Pred. No. 4.6e+05;
                                                                     Mismatches
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                                                                                                        DB 1;
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US-08-627-907A-6

Application US/08627907A

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HOLECULE TYPE: peptide

US-08-627-907A-6
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GENERAL INFORMATION:
APPLICANT: HIRANO, NAOTO
APPLICANT: HIRAI, Hisamaru
TITLE OF INVENTION: HUMAN PHOSPHOLIPASE C-ALPHA AND DNA
TITLE OF INVENTION: SEQUENCE ENCODING THE SAME
TITLE OF SECULENCES: 8
                                                                                                                                                                                                                                               Sequence 3, Applic Patent No. 6291205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
CURRENT APPLICATION NUMBER: US/07/901,713A
                    TITLE OF INVENTION: METHOD FOR INCREASING PRODUCTION OF TITLE OF INVENTION: DISULFIDE BONDED RECOMBINANT PROTEINS BY SACCAROMYCES TITLE OF INVENTION: CEREVISIAE FILE REFERENCE: 18469
                                                                                                APPLICANT:
                                                                                                                       APPLICANT:
                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (212) 596-9000
TELEPAX: (212) 596-9090
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                APPLICANT: Tuite, Michael F.
APPLICANT: Freedman, Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: HALEY Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: SHON-10
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMER: JP 5-238402
FILING DATE: 24-SEP-1993
PRIOR APPLICATION UNMER: PCT/JP94/01572
APPLICATION UNMER: PCT/JP94/01572
FILING DATE: 22-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS DOS SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: New York
STATE: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
                                                                                                                                                                                                                                                               Application US/07901713A
                                                                                            Montgomery, Donna L. Ellis, Ronald W.
                                                                                                                                                           Markus, Henry Z.
                                                                                                                                        Schultz, Loren D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: c/o FISH & NEAVE
1251 Avenue of the Americas
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100.0%; Pred. No. 4.6e+05;
htive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 6;
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US-08-750-142B-54
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                                                                 US-08-750-142B-54
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 Matches
               Query Match
Best Local (
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NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                            TELEPHONE: 617-542-50
TELEFAX: 617-542-8906
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 06:
TELECOMMUNICATION INFORMATION:
TRELEBHONE. 77
                                                                                                                                                                                                                                                      FILING DATE: 24-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Fraser Tanin "
                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08//50,112.

PILING DATE: 25-NOV-1996

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sarnstrand, TITLE OF INVENTION: NE
                                                                                MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: STREET: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM CON OPERATING SYSTEM:
                                                                                             TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                         NAME: Fraser, Janis K. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/
FILING DATE: 22-MAR-1996
APPLICATION NUMBER: SE95
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Similarity 6; Conserv
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 Conservative
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Lindvall, Magnus
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                  100.0%;
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               Score 47; DB 2; Length 12; Pred. No. 0.75;
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Best Local Similarity
6; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 5210073
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 105 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 1
CURRENT APPLICATION DATA:
                                                                                                                                                                          NAME: Billings, Lucy J.
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-
TELECOMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                  FILING DATE: Herewit CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Diskett
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APPLICANT: Shah, Purvi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 4
                               MMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 3174 FOR CITY: Palo Alto
LIBRARY: GenBank CLONE: 339649
                                                STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 28-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                TYPE: amino acid
                                                                                                                                                                TELEFAX: 415-845-4166
                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94304
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                                                                                               105 amino acids
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Best Local Similarity
Matches' 6; Conserv
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/826,910 FILING DATE: Herewith
                                                                                                                                                                                                                                                                                                NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                               IMMEDIATE SOURCE:
LIBRARY: GenBank
                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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APPLICANT: Shah, Pu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                    MBULL.
LIBRARY: GELL
ONE: 453972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT: Shah, Purvi
ITLE OF INVENTION: TH
                                                                                                                                                                                       LENGTH: 105 amino TYPE: amino acid STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
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ZIP: 94304
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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STREET: 31
                                                                                                                                                                            TOPOLOGY:
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31 WCGPCK 36
                          1 WCGPCK 6
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3174 Porter Drive
                                                                                                                                                                                                                        105 amino acids
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                                                        Conservative
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100.0%; Pred. No. 4.4;
                                                                       100.0%; Score 47; DB: 100.0%; Pred. No. 4.4;
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                                                         Mismatches
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RESULT 8 US-09-601-144-68

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Sequence 68, Application US/09601144 Patent No. 6566514

GENERAL INFORMATION:
APPLICANT: Wright, Jim A.
APPLICANT: Young, Aiping H.

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Query Match
Best Local Similarity
Thes 6; Conserv
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; ORGANISM: Human
US-09-601-144-68
                                                                                                                                                                                                                         US-08-386-729A-10
                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Homo sapiens US-09-513-999C-8037
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 74
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 68
                                                                                                                                                   Patent No. 5753435
GENERAL INFORMATION:
APPLICANT: Aharon
                                                                                                                                                                                         Sequence 10, Application US/08386729A Patent No. 5753435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 8037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PAIENT OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins Patent No. 6783961
FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                Best
                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lee, Yoon S.

TITLE OF INVENTION: OLIGONUCLEOTIDE SEQUENCES COMPLEMENTARY TO THIOREDOXIN TITLE OF INVENTION: AND THIOREDOXIN REDUCTASE GENES AND METHODS OF USING TITLE OF INVENTION: SAME TO MODULATE CELL GROWTH

FILE REFERENCE: 683-112US-A
CURRENT FILINGIATION NUMBER: US/09/601,144

CURRENT FILING DATE: 2000-10-18

PRIOR APPLICATION NUMBER: US 60/073,196

PRIOR PILING DATE: 1998-01-30

PRIOR FILING DATE: 1998-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 105
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                                                                                                     APPLICANT:
                                                     APPLICANT:
                                                                     APPLICANT:
                                                                                     APPLICANT:
                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31 WCGPCK 36
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                 Argaman, Anat
Av-Gay, Yossef
Nan, Helena M.
Kattevilder, Alfred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09513999C
                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                              Bovenberg, Roelor
Schreiber, Rachel
                                                                                                                                   Van Der Voort, Lucia H. M.
                                                                                                                   Cohen, Gerald
Pallissa, Harriet
                                                                                                                                                       Aharonowitz, Yair
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                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                     Roelof A. L.
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                                                                                                                                                                                                                                                                                                                                                                             Score 47; DB Pred. No. 4.4;
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Pred. No. 4.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                             DB 2; Length 105;
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RESULT 11
US-07-745-382-22
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Patent No. 527018:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (415) 854-3713
INFORMATION FOR SEQ ID NO: 10:
SECONDARY OF THE 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/274,043
FILING DATE: 12-JULY-1994
APPLICATION NUMBER: US 07/820,688
FILING DATE: 24-MARCH-1992
APPLICATION NUMBER: PCT/NL91/000101
FILING DATE: 18-JUNE-1991
FILING DATE: 18-JUNE-1991
FILING DATE: 18-JUNE-1991
                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: MCCOy, John
APPLICANT: LAVAllie, Edward
TITLE OF INVENTION: Peptide and
TITLE OF INVENTION: Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 18-JUNE-199
ATTORNEY/AGENT INFORMATION:
NAME: Rae-Venter, Barbar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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CITY: Menlo Park
STATE: CA
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                                                     COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
TOPOLOGY: linear
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                                                                                                           STATE:
                                                                                                                                                  CITY:
                                                                                                                                                                                 STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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                                      02140
                                                                                                                                              Cambridge
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                                                                                                        Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/07745382
                                                                                                                                                                     E: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            106 amino acids
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2882 Sand Hill Road, Ste. 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           An oxido reductase enzyme system obtainable from P. chrysogenum, the set of genes encoding the same and the use of oxido reductase enzyme systems or genes encoding the same for increasing antibiotic production
                                                                                                                                                                                                                                                                                                24
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CURRENT APPLICATION DATA:

PatentIn Release #1.0, Version #1.25

FILING DATE: APPLICATION NUMBER:

19910814

US/07/745,382

OPERATING SYSTEM:

IBM

Floppy disk

M PC compatible

TEM: PC-DOS/MS-DOS

MEDIUM TYPE:

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US-07-921-848-22
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ATTORNEY/AGENT INFORMATION:
NAME: CSERY, LUANN
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI5188A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-851
INFORMATION FOR SEQ ID NO: 22:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Lil
                                                                             FILING DATE: 06-FEB-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/745,382
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
NUMBER: CEST INFORMATION:
                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of COMPUTER: IBM PC comp
                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/652,531
                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                      TORNEL, COURT LUANN
NAME: CSET, LUANN
REGISTRATION NUMBER: 31,822
REGISTRATION NUMBER: G15188A
                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                APPLICATION NUMBER: US
FILING DATE: 19920728
             REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              109 amino acids
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(617) 876-1170
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Pred. No.
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RESULT 14 US-08-165-301A-26

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US-08-165-301A-22
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                                                                                                             US-08-165-301A-22
                                                                    Query Match
Best Local !
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                           TELEFAX: (617) 876-58
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTMARE: Patentin Palanaments
                                                     Local Similarity
nes 6; Conserv
                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE:
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32 WCGPCK 37
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                           1 WCGPCK 6
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TO NO: 22:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DiBlasio-Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  McCoy, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Grant
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876: 22:
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                                                                  100.0%;
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PEPTIDE AND PROTEIN FUSIONS TO
THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
THIOREDOXIN-LIKE MOLECULES
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Pred. No.
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                                                       Mismatches
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INFORMATION:

John

Application US/08165301A

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                                                                                                                                                                                                                                                                                                                                                                  Sequence 28, Application US/08165301A Patent No. 5646016
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                  GENERAL INFORMATION:

APPLICANT: McCoy, John
APPLICANT: Diblasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI 5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617) 876-58
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: DiBlasio-Smith, Eli
APPLICANT: Grant, Kathleen
APPLICANT: LaValle, Edward R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                              NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Cambridge
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                                                           COUNTRY:
                                                                                                   CITY: Cambridge
                                                                                                                        STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
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Y: U.S.A.
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                                                                                                                    E: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 47; DB 1; Length 109; milarity 100.0%; Pred. No. 4.6; Conservative 0; Mismatches 0: Tndele
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                                                           U.S.A.
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                                                                         Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                        TELEFAX: (617) 876-5851 INFORMATION FOR SEQ ID NO: 28:
                                                                                                                                                                                                                                                                                                            NAME: Meinert, M. C.
REGISTATION NUMBER: 33,544
REFERENCI/OCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
32 WCGPCK 37
                                      1 WCGPCK 6
                                                                             100.0%; Score 47; ilarity 100.0%; Pred. No. Conservative 0; Mismatch
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                                                                                                                 DB 1; Length 109;
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Search completed: February 23, 2006, 23:17:29 Job time : 48 secs

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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seq length: 2000000000
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Match Length
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15.286 Million cell updates/sec
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2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

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6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd.
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US-10-032-201B-105
US-10-032-201B-126
US-10-032-201B-128
US-10-501-282-3126
US-10-501-282-3126
US-10-032-513-7785
US-10-660-118A-7
US-10-451-467A-450
US-10-451-467A-450
US-10-451-467A-614
US-09-954-342-63
US-09-954-342-63
                                                                                                                                                                                                        US-09-801-260-4

US-10-145-586-43

US-10-425-115-351019

US-10-408-765A-312

US-10-424-599-280571

US-10-032-201B-143
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0-032-201B-102
0-032-201B-104
0-032-201B-105
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                    128, App
3126, Ap
96, Appl
7785, Ap
7785, Ap
77, Appl
27, Appl
450, App
614, App
62, Appl
63, Appl
63, Appl
64, Appl
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312, App
280571,
143, App
104, App
104, App
105, App
126, App
126, App
128, App
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SEQ ID NO 4

PRIOR APPLICATION NUMBER: US 60/333,866
PRIOR FILING DATE: 2001-11-28
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Microsoft Word

47 100.0 104 4 US-10-032-201B-154 47 100.0 104 4 US-10-032-201B-154 47 100.0 104 4 US-10-032-201B-158 47 100.0 104 4 US-10-032-201B-158 47 100.0 104 4 US-10-032-201B-159 47 100.0 104 4 US-10-032-201B-161 47 100.0 104 4 US-10-032-201B-161 47 100.0 104 4 US-10-032-201B-276 47 100.0 104 4 US-10-424-599-28258 47 100.0 104 4 US-10-424-599-28258 47 100.0 105 3 US-10-451-467A-188 47 100.0 105 4 US-10-316-253-291 47 100.0 105 4 US-10-316-253-291 47 100.0 105 4 US-10-312-201B-106 47 100.0 105 4 US-10-312-201B-106 47 100.0 105 4 US-10-32-201B-106 47 100.0 105 4 US-10-32-201B-107 47 100.0 105 4 US-10-351-891-127 47 100.0 105 4 US-10-351-891-127 47 100.0 105 4 US-10-351-891-127 47 100.0 105 4 US-10-660-118A-8	47 100.0 104 4 47 100.0 104 4 47 100.0 104 4 47 100.0 104 4 47 100.0 104 4 47 100.0 104 4 47 100.0 104 4 47 100.0 105 3 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4 47 100.0 105 4
100.0 10	100.0 104 4 US-10-032-201B-154 Sequence 100.0 104 4 US-10-032-201B-154 Sequence 100.0 104 4 US-10-032-201B-158 Sequence 100.0 104 4 US-10-032-201B-158 Sequence 100.0 104 4 US-10-032-201B-159 Sequence 100.0 104 4 US-10-032-201B-161 Sequence 100.0 104 4 US-10-032-201B-161 Sequence 100.0 104 4 US-10-032-201B-258 Sequence 100.0 104 4 US-10-424-592-28588 Sequence 100.0 104 4 US-10-451-467A-188 Sequence 100.0 105 4 US-10-171-311-226 Sequence 100.0 105 4 US-10-316-253-291 Sequence 100.0 105 4 US-10-032-201B-16 100.0 105 4 US-10-660-118A-8
	104 4 US-10-032-201B-152 Sequence 104 4 US-10-032-201B-154 Sequence 104 4 US-10-032-201B-158 Sequence 104 4 US-10-032-201B-159 Sequence 104 4 US-10-032-201B-159 Sequence 104 4 US-10-032-201B-161 Sequence 104 4 US-10-032-201B-161 Sequence 104 4 US-10-032-201B-161 Sequence 104 4 US-10-032-201B-276 Sequence 105 4 US-10-451-467A-188 Sequence 105 4 US-10-451-467A-188 Sequence 105 4 US-10-171-311-226 Sequence 105 4 US-10-316-253-291 Sequence 105 4 US-10-032-201B-10 105 4 US-10-040-765A-2019 105 4 US-10-060-11BA-8
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4 US-10-032-201B-154 4 US-10-032-201B-158 4 US-10-032-201B-158 4 US-10-032-201B-159 4 US-10-032-201B-161 4 US-10-032-201B-164 6 US-10-032-201B-164 6 US-10-424-599-28258 6 US-10-451-467A-188 7 US-10-451-467A-188 7 US-10-171-311-226 7 US-10-316-253-291 7 US-10-032-201B-106	-10-032-2018-152 -10-032-2018-153 -10-032-2018-155 -10-032-2018-155 -10-032-2018-159 -10-032-2018-159 -10-032-2018-161 -10-032-2018-161 -10-032-2018-161 -10-032-2018-161 -10-032-2018-161 -10-032-2018-161 -10-424-599-282588 -10-451-467A-188 -10-451-467A-188 -10-311-226 -10-171-311-226 -10-316-253-291 -10-32-2018-10 -10-3
US-10-032-201B-154 US-10-032-201B-154 US-10-032-201B-158 US-10-032-201B-158 US-10-032-201B-159 US-10-032-201B-161 US-10-032-201B-276 US-10-032-201B-276 US-10-451-467A-188 US-10-451-47A-188 US-10-451-47A-188 US-10-451-47A-188 US-10-451-47A-188 US-10-451-47A-188 US-10-451-47A-188 US-10-451-47A-188 US-10-451-47A-188 US-10-451-311-226 US-10-118-23-291 US-10-032-201B-106 US-10-032-201B-106 US-10-032-201B-107 US-10-404-599-237784 US-10-408-758A-2019 US-10-4060-118A-8	-10-032-2018-152 -10-032-2018-153 -10-032-2018-155 -10-032-2018-155 -10-032-2018-159 -10-032-2018-159 -10-032-2018-161 -10-032-2018-161 -10-032-2018-161 -10-032-2018-161 -10-032-2018-161 -10-032-2018-161 -10-424-599-282588 -10-451-467A-188 -10-451-467A-188 -10-311-226 -10-171-311-226 -10-316-253-291 -10-32-2018-10 -10-3
	sequence

ALIGNMENTS

RESULT 1 US-10-304-287-4

GENERAL INFORMATION:
APPLICANT: Waisman, David M.
APPLICANT: Kwon, Mijung

APPLICANT: Kwon, Mijung TITLE OF INVENTION: Anti-angiogenesis methods, compositions and uses therefor FILE REFERENCE: ME02-001
CURRENT APPLICATION NUMBER: US/10/304,287
CURRENT FILING DATE: 2002-11-26

Sequence 4, Application US/10304287 Publication No. US20030083234A1

```
Sequence 3, Application US/10660118A

Publication No. US20040131606A1

GENERAL INFORMATION:

APPLICANT: White, Carl W.

ITILE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum FILE REFERENCE: 2879-98

CURRENT APPLICATION NUMBER: US/10/660,118A

CURRENT APPLICATION NUMBER: US/10/660,118A

CURRENT FILING DATE: 2003-09-10

PRIOR APPLICATION NUMBER: 60/409,960

PRIOR FILING DATE: 2002-09-10

PRIOR FILING DATE: 2003-04-11

NUMBER OF SEQ ID NOS: 15

SOFTWARE: PatentIn version 3.1

INDUSTRIANS OF SEQ ID NOS 3.1
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Matches 6; Conser
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                    LENGTH: 6
TYPE: PRT
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TYPE: PRT
ORGANISM: mammalian
ORGANISM: Artificial sequence
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Sequence 4, Application US/09801260

Patent No. US20020034801A1

GENERAL INFORMATION:
APPLICANT: Curtis, Rory A.J.
TITLE OF INVENTION: 22105, A NOVEL HUMAN THIOREDOXIN FAMILY
TITLE OF INVENTION: 22001

CURRENT APPLICATION NUMBER: US/09/801,260

CURRENT APPLICATION NUMBER: US/09/801,260

CURRENT FILING DATE: 2001-03-06

PRIOR APPLICATION NUMBER: US 60/187,447

PRIOR FILING DATE: 2000-03-07

NUMBER: OF SEQ ID NOS: 5

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4

LENGTH: 52
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US-10-735-577-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; FEATURE: ; OTHER INFORMATION: synthetic peptide motif US-10-660-118A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-801-260-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-735-577-4
                                                                                                                                               US-09-801-260-4
                                                                                      Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Microsoft Word
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/10735577 Publication No. US20040142897A1 GENERAL INFORMATION:
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                                                                                                           Query Match
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TITLE OF INVENTION: Compositions and Methods for Inhibiting Tumor Growth and Metastas
FILE REFERENCE: ME03-009
CURRENT APPLICATION NUMBER: US/10/735,577
CURRENT FILING DATE: 2003-12-12
PRIOR APPLICATION NUMBER: US 60/433,140
PRIOR FILING DATE: 2002-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2002-12-13
NUMBER OF SEQ ID NOS: 160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 6
TYPE: PRT
                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: mammalian
                                                                                                                                                               OTHER INFORMATION: consensus sequence
                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity nes 6; Conserv
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les 6; Conserv
                                                                                        Local
 20 WCGPCK 25
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                                                                                        Similarity
                                     WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 47; DB 4; 1 ilarity 100.0%; Pred. No. 1.7e+06; Conservative 0; Mismatches 0;
                                                                   100.0%; ilarity 100.0%; Conservative 0
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                                                                         0
                                                                       Score 47; DB Pred. No. 7.1; Mismatches
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                                                                                                         DB 3; Length 52;
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                                                                       Gaps
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RESULT 7

US-10-408-765A-312

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; OTHER INFORMATION: Clone ID: MRT4577_83299C.1.pep
US-10-425-115-351019
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 43, Application US/10145586
Publication No. US20030138890A1
                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 351019
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                                                                                 Matches
                                                                                                                      Query Match
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Best Local
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APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: M. Galvin, Katherine
                                                                                                Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
APPLICANT:
                                                                                                                                                                                                        LENGTH: 82
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Nucleic Acid Molecules
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: La Rosa, Thomas
APPLICANT: Kovalic, David
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 95
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Bandaru, Rajasekhar
APPLICANT: Kapeller-Libermann, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS,
TITLE OF INVENTION: HUMAN THOREDOXIN FAMILY MEMBERS, HUMAN LEUCINE-RICH
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBER
FILE REFERENCE: 10448-188001
CURRENT APPLICATION NUMBER: US/10/145,586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT FILING DATE: 2002-05-14
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ORGANISM: Artificial Sequence
FEATURE:
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les 6; Conserv
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5 WCGPCK 10
                                       1 WCGPCK 6
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                                                                               Conservative
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                                                                                                     100.0%; Score 47; DB 100.0%; Pred. No. 10;
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                                                                                   Mismatches
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                                                                                                                        DB 4; Length 82
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RESULT 9
US-10-032-201B-143
; Sequence 143, Application US/10032201B
; Publication No. US20030167524A1
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Best Local S
Matches 6
GENERAL INFORMATION
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SEQ ID NO 312
LENGTH: 84
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 280571
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Publication No. US20040031072A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
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                                                                                                                                                                                                                                                                            FEATURE:
OTHER INFORMATION: Clone ID: PAT_MRT3847_95379C.1.pep
10-424-599-280571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/408,765A CURRENT FILING DATE: 2003-04-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 3077
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ITTLE OF INVENTION: TARGETS FOR THERAPEUTIC INTERVENTION

ITTLE OF INVENTION: IDENTIFIED IN THE MITOCHONDRIAL PROTEOME
                                                                                                                                                                                                                                                                                                                                 NAME/KEY: unsure
LOCATION: (1)..(90)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 90
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                                                                                                                              13 WCGPCK 18
                                                                                                                                                               1 WCGPCK 6
                                                                                                                                                                                                 l Similarity
6; Conserv
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Taylor, Steven W.
Glenn, Gary M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zhou Yihua
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                                                                                                                                                                                                                      Score 47; DB 4;
Pred. No. 11;
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                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                          Length 90;
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                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 102
                                                              Matches
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                                                                             Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/032,2018
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                        FILE REPERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032;201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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ORGANISM: Thiocapsa roseopersicina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT
                                                                                                                                         TYPE: PRT
ORGANISM: Chlamydia muridarum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH: 91
                                                                                                                                                                            ENGTH: 102
                                                                                                                                                                                                                                                                                          TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      102, Application US/10032201B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     32
                                                                                                                                                                                                                                                                                        I: Moloney, Maurice INVENTION: METHODS FOR THE PRODUCTION INVENTION: COMPOSITIONS
                                                            9
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WCGPCK 32
                               WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                        Heifetz, Peter Bernard
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zaplachinski, Steve
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ilarity 100.0%;
Conservative 0
                                                               Conservative
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                                                                           100.0%;
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                                                              Score 47; DB Pred. No. 12; 0; Mismatches
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Pred. No. 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                          OF MULTIMERIC PROTEINS,
                                                                                          Length 102;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
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                                                                                                                                                                                                                                                                                                            AND RELATED
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RESULT 11
US-10-032-201B-104
; Sequence 104; Application US/10032201B
; Publication NO. US20030167524A1

GENERAL

INFORMATION:

APPLICANT:

Van Rooijen, Gijs

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Sequence 126, Application US/10032201B
Publication No. US20030167524A1
GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 105
LENGTH: 102
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CURRENT APPLICATION NUMBER: US/10/032,2018
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 104
LENGTH: 102
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Best Local
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-10-032-201B-104
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
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Heifetz, Peter Bernard
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Zaplachinski,
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Dalmia, Bipin Kumar
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Dalmia, Bipin Kumar
Briggs, Steven
Dalmia, Bipin Kumar
Del Val, Greg
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Zaplachinski, Steve
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b. US20030167524A1
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 126
SEQ ID NO 126
LENGTH: 102
                                                                                                                                            Sequence 3126, Application US/10501282
Publication No. US20050203280A1
GENERAL INFORMATION:
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SEQ ID NO 128
LENGTH: 102
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APPLICANT:
APPLICANT:
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
APPLICANT: MCMICHAEL, JOHN CALHOUN
APPLICANT: ZAGURSKY, ROBERT JOHN
APPLICANT: RUSSELL, DAVID PARRISH
APPLICANT: FLETCHER, LEAH DIANE
TITLE OF INVENTION: ALLOICOCCUS OTITIDIS OPEN READING FRAMES (ORFS) ENCODING
TITLE OF INVENTION: POLYPEPTIDE ANTIGENS, IMMUNOGENIC COMPOSITIONS AND USES 7
FILE REFERENCE: AM100780 L2
CURRENT APPLICATION NUMBER: US/10/501,282
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter B
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
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Heifetz, Peter Bernard
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Zaplachinski, Steve
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Dalmia, Bipin Kumar
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Pred. No.
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PRIOR APPLICATION NUMBER: 60/333,777

PRIOR FILING DATE: 2001-11-29

PRIOR APPLICATION NUMBER: 60/426,742

PRIOR APPLICATION NUMBER: 60/426,742

PRIOR APPLICATION NUMBER: PCT/US02/36123

PRIOR APPLICATION NUMBER: 60/202/36123

PRIOR PILING DATE: 2002-11-25

NUMBER OF SEQ ID NOS: 6653

SOFTWARE: Patentin version 3.2

SEQ ID NO 3126

PRIOR APPLICATION NUMBER: 60/202/36123

SEQ ID NO 3126

PRIOR PILING DATE: 2002-11-18

PRIOR APPLICATION NUMBER: 60/202/36123

SEQ ID NO 3126

PRIOR APPLICATION NUMBER: 60/202/36123

SOFTWARE: Patentin version 3.2

SOFTWARE: Patentin version 3.2

SEQ ID NO 3126

PRIOR APPLICATION NUMBER: 66: 00/202/36123

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SEQ ID NO 3126

PRIOR APPLICATION NUMBER: 66: 00/202/36123

SOFTWARE: Patentin version 3.2

SEQ ID NO 3126

PRIOR APPLICATION NUMBER: pct/US02/36123

SOFTWARE: Patentin version 3.2

SEQ ID NO 3126

PRIOR APPLICATION NUMBER: pct/US02/36123

SOFTWARE: Patentin version 3.2

PRIOR APPLICATION NUMBER: pct/US02/36123

SOFTWARE: patentin version 3.2

SOFTWARE: patentin version 3.
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Sequence Sequence Sequence

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12, Appl 16, Appl 10, Appl 11, Appl 13, Appl 14, Appl 15, Appl 15, Appl 15, Appl 8494, App 4703, App 4703, Appl 470, Appl 60, Appl 114, Appl 114, Appl 117, Appl 118, Appl 118, Appl 118, Appl 118, Appl

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Total number of hits satisfying chosen parameters:
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1: /cgn2 6/ptodata/1/pubpaa/US08 NEW PUB.pep:*

2: /cgn2 6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

3: /cgn2 6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

4: /cgn2 6/ptodata/1/pubpaa/PCT NEW PUB.pep:*

5: /cgn2 6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

6: /cgn2 6/ptodata/1/pubpaa/US01 NEW PUB.pep:*

7: /cgn2 6/ptodata/1/pubpaa/US1 NEW PUB.pep:*

8: /cgn2 6/ptodata/1/pubpaa/US1 NEW PUB.pep:*
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Maximum Match 100%
Listing first 45 summaries
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US-11-090-916-8

US-11-090-916-9

US-11-090-916-10

US-11-090-916-11

US-11-090-916-12

US-11-090-916-13

US-11-090-916-13

US-11-144-301A-9

US-11-144-301A-10

US-11-090-916-4

US-11-090-916-4

US-11-090-916-4

US-11-106-796-11

US-11-106-796-11

US-11-106-796-11

US-11-106-796-11

US-11-131-744-3

US-11-131-744-6

US-11-131-744-7

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US-11-090-916-3
Sequence 3, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: White, Carl W.
APPLICANT: Lee, Rees
ITITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/550,865
PRIOR FILING DATE: 2003-09-0
INUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.3
SEQ ID NO 3
LENGTH: 6
TUDE: DBT
  RESULT 2
US-11-131-744-2
US-11-131-744-2
; Sequence 2, Application US/11131744
; Publication No. US20060023808A1
; GENERAL INFORMATION:
; APPLICANT: HAJIVANDI, MAHBOD R.
; APPLICANT: POPE, ROBERT M.
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Best Local Similarity
Matches 6; Conserv
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ORGANISM: Artificial sequence
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US-11-090-916-13
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US-10-467-657-8494
US-10-962-561-703
US-10-131-826A-426
US-10-821-234-860
US-10-821-234-860
US-10-821-234-287
US-10-821-234-1287
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US-10-821-234-1287
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Pred. No. 8.3e+04
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length: 0 length: 2000000000

Title: Perfect score:

US-10-660-118A-3

February 23,

Copyright

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Sequence:

Scoring table:

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CURRENT FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/572,290
PRIOR PELLING DATE: 2005-05-17
PRIOR PELLING DATE: 2005-05-17
PRIOR PELLING DATE: 2005-04-08
PRIOR FILING DATE: 2005-04-08
PRIOR PELLING DATE: 2004-05-17
PRIOR PELLING DATE: 2004-05-17
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR PILING DATE: 2004-10-26
PRIOR PILING DATE: 2004-10-26
PRIOR PILING DATE: 2004-10-26
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Best Local Similarity
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LENGTH: 75
                                                                                                                                                     NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver.
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/11131744
Publication No. US20060023808A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR CALIBRATION IN MASS TITLE OF INVENTION: SPECTROMETRY FILE REFERENCE: NVX-28 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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SOFTWARE: PatentIn Ver. 3.3
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PRIOR APPLICATION NUMBER: 60/572,215
PBIOR FILING DATE: 2004-05-17
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CURRENT FILING DATE: 2005-05-17
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                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/621,686 PRIOR FILING DATE: 2004-10-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIGR APPLICATION NUMBER: 60/669,373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
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ORGANISM: Artificial Sequence
OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
                                                    ORGANISM: Artificial Sequence FEATURE:
                                                                                                                             ENGTH: 91
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LEITE, JOHN F
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LEITE, JOHN F.
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; ORGANISM: Saccharomyces cerevisiae US-11-090-916-7
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NUMBER OF SEQ ID NOS: 381
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 286
LENGTH: 103
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Best Local Similarity
Matches 6; Conserv
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                                                                                      SEQ ID NO 7
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Best Local Similarity
                                                                                                                                           TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-09-10
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
PRIOR PILING DATE: 2005-02-07
PRIOR FILING DATE: 2005-02-07
PRIOR FILING DATE: 2005-02-07
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APPLICANT: McAuliffe, Olivia
APPLICANT: Peril, Andrea Azcarate
TITLE OF INVENTION: Nucleic Acid Sequences Encoding
TITLE OF INVENTION: Stress-Related Proteins and Uses Therefore
FILE REFERENCE: 5051-694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/11/074,176
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: 60/551,161
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                                                                                                         NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn versi
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: White, Carl W. APPLICANT: del Val, Greg APPLICANT: Lee, Rees
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                                           TYPE: PRT
                                                             ENGTH: 103
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100.0%; Pred. No. 0.23;
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100.0%; Pred. No.
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SEQ ID NO 137
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                                                                                          SEQ ID NO 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-821-234-1371
                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10
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TITLE OF INJENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
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                                            ORGANISM: Gallus gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                          ENGTH:
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                                                                                                                                                                                                             FILING DATE: 2003-04-11
APPLICATION NUMBER: 10/660,118
                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/462,082
                                                                                                                                                                          FILING DATE: 2003-09-10
APPLICATION NUMBER: 60/556,516
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                                                                                                                                                                     LING DATE:
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                                                                                                        PatentIn version 3.3
                                                                                                                                                     ICATION NUMBER: 60/650,865
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 Score 47;
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Length 105;
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SEQ ID NO 9
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Publication No.
NUMBER OF
                                PRIOR FILING DATE: 2004-0 PRIOR APPLICATION NUMBER:
                                                  PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
                                                                                                                                                                                                                                                                                          APPLICANT: White, Car
APPLICANT: del Val,
                 PRIOR FILING DATE:
                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10
                                                                                                                                                                                                   CURRENT FILING DATE: 2005-03-24
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CURRENT APPLICATION NUMBER: US/11/090,916
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CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
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ORGANISM: Mus musculus
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2005-02-07
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                                60/650,865
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US-11-090-916-11
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Publication No. US20050260140A1
GENERAL INFORMATION:
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LENGTH: 105
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                                                                                                FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 66/462,082
PRIOR APPLICATION NUMBER: 66/462,082
PRIOR FILING DATE: 2003-04-11
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CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
                                   PRIOR APPLICATION NUMBER: 10/660,118 PRIOR FILING DATE: 2003-09-10 PRIOR APPLICATION NUMBER: 60/556,516
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TITLE OF INVENTION: Mucclytic and Anti-Elastase Compounds and Methods of Use
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PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
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les 6; Conserv
APPLICATION NUMBER: 60/650,865
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; PRIOR FILING DATE: 2005-02-07; NUMBER OF SEQ ID NOS: 15; SOFTWARE: PatentIn version 3.3; SEQ ID NO 12; LENGTH: 105; TYPE: PRT; ORGANISM: Homo sapiens US-11-090-916-12
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US-11-144-301A-9
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; ORGANISM: Homo sapiens
US-11-144-301A-9
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                                                                                                                             PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
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CURRENT APPLICATION NUMBER: US/11/144,301A
CURRENT FILING DATE: 2005-06-03
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PRIOR FILLING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
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APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
                                                                                         NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                   APPLICANT: Johanna, Ungerstedt
TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
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                                     LENGTH: 10
TYPE: PRT
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ORGANISM: Homo sapiens
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RESULT 14
US-11-144-301A-10
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; SEQ ID NO 10
; LENGTH: 105
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-144-301A-10
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Best Local Similarity 100.0%;
Matches 6; Conservative 0
Query Match
Best Local Similarity
Matches 6; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 943
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                                                                                               LENGTH: 105
TYPE: PRT
ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank BAA04881
DATABASE ENTRY DATE: 2002-12-25
:-11-032-773-943
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Publication No. US20050288227A1
GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/11/032,773
CURRENT FILING DATE: 2005-01-11
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PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
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TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
FILE REFERENCE: 28517-501 UTIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Marks, Paul A. APPLICANT: Johanna, Unge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Levy, Ronald
TITLE OF INVENTION: DESIGN OF THERAPEUTICS AND THERAPEUTICS
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APPLICANT: Levy, Ronald
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31 WCGPCK 36
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    100.0%; Score 47; DB 7; Length 105; ilarity 100.0%; Pred. No. 0.25; Conservative 0; Mismatches 0; Indels
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